

GenCore version 5.1.3
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OM protein - protein search, using sw model

Run on: December 30, 2002, 16:15:48 ; Search time 17 Seconds
(without alignments)
367.573 Million cell updates/sec

Title: US-09-664-326-23

Perfect score: 368
Sequence: 1 LTYVDCYTESGQNLCLCEGSSN.....PKQSHNDGFEEIPEYLQ 65

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283224 seqs, 96134422 residues

Total number of hits satisfying chosen parameters: 283224

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :
1: PIR1:*
2: PIR2:*
3: PIR3:*
4: PIR4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	360	97.8	65	1 HULXH	thrombin inhibitor
2	353	95.9	65	2 S05674	hirudin IIB - medi
3	352	95.7	65	2 S78521	hirudin IIB' - me
4	350	95.1	65	2 S05673	hirudin IIA - medi
5	348	94.6	65	2 S05676	hirudin IIA' - medi
6	346	94.0	65	2 S05678	hirudin IIA' - medic
7	345	93.8	65	2 S78520	hirudin IIA' - me
8	345	93.8	65	2 S05672	hirudin IIB - med
9	345	93.8	65	2 S05675	hirudin IIB - med
10	343	93.2	65	2 S05679	hirudin IIB' - medi
11	333	90.5	72	2 A37417	thrombin inhibitor
12	315	85.6	66	2 A24350	thrombin inhibitor
13	288	78.3	55	2 S05672	hirudin I - medic
14	249	67.7	63	2 A53883	hirudin HVI homolo
15	247	67.1	84	2 S33329	hirudin HMI - leec
16	230	62.5	84	2 S33328	hirudin HMI - leec
17	215	58.4	63	1 A42207	hirudin p6 - leech
18	173	47.0	62	1 HULXH	hirudin p18 - lee
19	90	24.5	17	2 S05671	hirudin Ia - medic
20	74	20.1	2352	2 T30201	Notch homolog prot
21	69	18.8	21	2 A42125	trophozoite cystel
22	68.5	18.6	1071	1 PXYVVA	H+-exporting ATPas
23	68.5	18.6	2233	2 T28669	surface protein 51
24	66	17.9	761	2 T09052	hypothetical prote
25	66	17.9	2150	2 T32497	hypothetical prote
26	66	17.9	5376	2 T42215	zonahesin - mouse
27	65	17.7	558	2 T15448	hypothetical prote
28	65	17.7	1043	2 T19734	hypothetical prote
29	65	17.7	1661	2 T31330	head-activator bin

30	64	17.4	317	2 I46916	insulin-like growth
31	64	17.4	1743	2 T26859	hypothetical prote
32	63.5	17.3	318	2 A82319	glutathione synthe
33	63.5	17.3	474	2 S18452	variant surface gl
34	63.5	17.3	755	2 A44315	collagen oligomer
35	63.5	17.3	1722	2 E89753	protein F11C7.4 [l
36	63.5	17.3	2321	2 S78549	notch3 protein - h
37	63.5	17.3	2703	1 A24420	antifungal protein
38	63	17.1	80	2 T10183	S-receptor kinase
39	63	17.1	778	2 T05341	activin receptor I
40	62.5	17.0	513	1 J01486	latent transformin
41	62.5	17.0	1820	2 A55494	antifungal protein
42	62	16.8	79	2 T10243	antifungal protein
43	62	16.8	79	2 T07917	probable antifunga
44	62	16.8	80	2 T02621	probable antifunga
45	62	16.8	80	2 T02622	probable antifunga

ALIGNMENTS

RESULT 1
HULXH
thrombin inhibitor (hirudin) - medicinal leech
C/Species: Hirudo medicinalis (medicinal leech)
C/Date: 30-Nov-1980 #sequence_revision 03-Aug-1984 #text_change 07-May-1999
C/Accession: A91318; A94429; A60811; A01289
R/Dodt, J.; Muller, H.P.; Seemuller, U.; Chang, J.Y.
FEBS Lett. 165, 180-183, 1984
A/Title: The complete amino acid sequence of hirudin, a thrombin specific inhibitor.
A/Reference number: A91318
A/Accession: A91318
A/Molecule type: protein
A/Residues: 1-65 <DOD>
R/Petersen, T.E.; Roberts, H.R.; Sottrup-Jensen, L.; Magnusson, S.; Bagdy, D.
In: Proteases of the Biological Fluids, Proc. 23rd Colloq., Peeters, H., ed., pp.145-14
A/Reference number: A94429
A/Accession: A94429
A/Molecule type: protein
A/Residues: 1-65 <P>
R/Mao, S.J.T.; Yates, M.T.; Blankenship, D.T.; Cardin, A.D.; Kristiansky, J.L.; Loven
Anal. Biochem. 161, 514-518, 1987
A/Title: Rapid purification and revised amino-terminal sequence of hirudin: a specific
A/Reference number: A60811; MUID:87211066; PMID:3578808
A/Accession: A60811
A/Molecule type: protein
A/Residues: 1-32, 'N', 34-43 <MAO>
A/Note: the authors suggest that their identification of 33-Asn is correct and that 3
present a natural variant of hirudin
C/Comment: Hirudin is a potent thrombin-specific protease inhibitor.
C/Superfamily: thrombin inhibitor
C/Keywords: anticoagulant; sulfoliprotein
F:6-14,16-28,22-39/Disulfide bonds: #status experimental
F:63/Binding site: sulfate (Tyr) (covalent) #status experimental

Query Match 97.8%; Score 360; DB 1; Length 65;
Best Local Similarity 96.9%; Pred. No. 7.8e-30;
Matches 63; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 LTYVDCYTESGQNLCLCEGSSNVCQGQNKCIIGSDGKNCQVYGEETPRQSHNDGFEEIP 60
DB 1 VVYVDCYTESGQNLCLCEGSSNVCQGQNKCIIGSDGKNCQVYGEETPRQSHNDGFEEIP 60
QY 61 EBYLQ 65
DB 61 EBYLQ 65

RESULT 2
S05674
hirudin IIB - medicinal leech
N/Alternate names: thrombin inhibitor
C/Species: Hirudo medicinalis (medicinal leech)

C>Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 17-Jul-1998
C:Accession: S05674
R:Scharf, M.; Engels, J.; Tripiier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-thirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S05674
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 95.9%; Score 353; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 4e-29;
Matches 61; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

OY 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60
DB 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60

OY 61 EYTLQ 65
DB 61 EYTLQ 65

RESULT 3
S05674
hirudin IIID' - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 17-Jul-1998
A:Accession: S78521
R:Scharf, M.; Engels, J.; Tripiier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-thirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S78521
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 95.7%; Score 352; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 5e-29;
Matches 61; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

OY 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60
DB 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60

OY 61 EYTLQ 65
DB 61 EYTLQ 65

RESULT 4
S05673
hirudin IIA - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
A:Accession: S05673
R:Scharf, M.; Engels, J.; Tripiier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-thirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Molecule type: protein
A:Residues: 1-65 <SCH>

C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:47/inhibitory site: lys (thrombin) #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 95.1%; Score 350; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 8e-29;
Matches 61; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60
DB 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60

OY 61 EYTLQ 65
DB 61 EYTLQ 65

RESULT 5
S05676
hirudin III - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
A:Accession: S05676
R:Scharf, M.; Engels, J.; Tripiier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-thirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S05676
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 94.6%; Score 348; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 1.3e-28;
Matches 61; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60
DB 1 LVYTCESGQMLCEGSSNVCGGKNCITLSDGKNCVTEGTPKPSHNDGDFEETP 60

OY 61 EYTLQ 65
DB 61 EYTLQ 65

RESULT 6
S05678
hirudin II - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 17-Jul-1998
A:Accession: S05678
R:Scharf, M.; Engels, J.; Tripiier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-thirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S05678
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 94.6%; Score 348; DB 2; Length 65;
Best Local Similarity 92.3%; Pred. No. 1.3e-28;
Matches 60; Conservative 5; Mismatches 0; Indels 0; Gaps 0;

QY 1 LVTYDCTESGONLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||
Db 1 IFTYDCTESGODLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
QY 61 EBYLQ 65
|||||
Db 61 EBYLQ 65

RESULT 7

578520
hirudin IIRa' - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C>Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 17-Jul-1998
C:Accession: S78520
R:Scharf, M.; Engels, J.; Tripler, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S78520
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match

Best Local Similarity 94.0%; Score 346; DB 2; Length 65;
Best Local Similarity 90.8%; Pred. No. 2e-28;
Matches 59; Conservative 5; Mismatches 1; Indels 0; Gaps 0;

QY 1 LVTYDCTESGONLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||
Db 1 VVTYDCTESGDLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
QY 61 EBYLQ 65
|||||
Db 61 EBYLQ 65

RESULT 8

505677
hirudin IIRb - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C>Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
C:Accession: S05677
R:Scharf, M.; Engels, J.; Tripler, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S05677
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match

Best Local Similarity 93.8%; Score 345; DB 2; Length 65;
Best Local Similarity 92.3%; Pred. No. 2.6e-28;
Matches 60; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 LVTYDCTESGONLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||
Db 1 VVTYDCTESGONLCLCGSDNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
QY 61 EBYLQ 65
|||||
Db 61 EBYLQ 65

RESULT 9

505675
hirudin IIRa' - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C>Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
C:Accession: S05675
R:Scharf, M.; Engels, J.; Tripler, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S05675
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match

Best Local Similarity 93.8%; Score 345; DB 2; Length 65;
Best Local Similarity 92.3%; Pred. No. 2.6e-28;
Matches 60; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 LVTYDCTESGONLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||
Db 1 VVTYDCTESGONLCLCGSDNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
QY 61 EBYLQ 65
|||||
Db 61 EBYLQ 65

RESULT 10

505679
hirudin IIR' - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C>Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 17-Jul-1998
C:Accession: S05679
R:Scharf, M.; Engels, J.; Tripler, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945; PMID:2792365
A:Accession: S05679
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match

Best Local Similarity 93.2%; Score 343; DB 2; Length 65;
Best Local Similarity 90.8%; Pred. No. 4.1e-28;
Matches 59; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

QY 1 LVTYDCTESGONLCLCGSNVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60
:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||
Db 1 IITYDCTESGODLCLCGSDVCGGKNCILGSDGKNCQVTGEGTPKPOSHNDGDFEIP 60

QY 61 EBYLQ 65
|||||
Db 61 EBYLQ 65

RESULT 11

A37417
thrombin inhibitor (hirudin) type II precursor - medicinal leech (fragment)
C:Species: Hirudo medicinalis (medicinal leech)
C>Date: 28-Apr-1993 #sequence_revision 28-Apr-1993 #text_change 16-Jul-1999
C:Accession: A37417
R:Harvey, R.P.; Degryse, E.; Stefani, L.; Schanber, F.; Cazenave, J.P.; Courtney, M.;
Proc. Natl. Acad. Sci. U.S.A. 83, 1084-1088, 1986

A:Title: Cloning and expression of a cDNA coding for the anticoagulant hirudin from the
A:Reference number: A37417; MUID:86149219; PMID:3531162
A:Accession: A37417
A:Status: preliminary; not compared with conceptual translation
A:Molecule type: mRNA
A:Residues: 1-72 <HAP>
A:Cross-references: GB:M12693; NID:g159224; PIDN:AAA29195.1; PID:g159225
C:Superfamily: thrombin inhibitor

Query Match 90.5%; Score 333; DB 2; Length 72;
Best Local Similarity 87.7%; Pred. No. 4.6e-27;
Matches 57; Conservative 6; Mismatches 2; Indels 0; Gaps 0;

QY 1 LVTDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 60
:|||||
DB 8 ITVDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 67
QY 61 EYIQQ 65
:|||||
DB 68 EYIQQ 72

RESULT 12

A24350
thrombin inhibitor (hirudin PA) - medicinal leech
N:Alternate names: hirudin (plasminogen activator-type)
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 18-Jun-1993
C:Accession: A24350
R:Doty, J.; Machalski, W.; Seemuller, U.; Maschler, R.; Fritz, H.
Biol. Chem. Hoppe-Seyler 367, 803-811, 1986
A:Title: Isolation and characterization of hirudin isolinhibitors and sequence analysis
A:Reference number: A24350; MUID:87026247; PMID:3768144
A:Accession: A24350
A:Molecule type: Protein
A:Residues: 1-66 <DDO>
C:Superfamily: thrombin inhibitor

Query Match 85.6%; Score 315; DB 2; Length 66;
Best Local Similarity 87.1%; Pred. No. 2.9e-25;
Matches 54; Conservative 4; Mismatches 4; Indels 0; Gaps 0;

QY 1 LVTDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 60
:|||||
DB 1 ITVDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 60
QY 61 EE 62
:|:
DB 61 ED 62

RESULT 13

S05672
hirudin I - medicinal leech (fragments)
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
C:Accession: S05672
R:Scharf, M.; Engels, J.; Tripler, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:9005945; PMID:2792365
A:Accession: S05672
A:Molecule type: Protein
A:Residues: 1-55 <SCB>
C:Superfamily: thrombin inhibitor
C:Key words: anticoagulant; serine proteinase inhibitor; sulfoprotein
F;6-14/Disulfide bonds: #status predicted
F;53/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 78.3%; Score 288; DB 2; Length 55;
Best Local Similarity 81.5%; Pred. No. 1.3e-22;
Matches 53; Conservative 1; Mismatches 1; Indels 10; Gaps 1;

QY 1 LVTDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 60
:|||||
DB 1 VYVDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 50
QY 61 EYIQQ 65
:|||||
DB 51 EYIQQ 55

RESULT 14

A53883
hirudin HVI homolog bufrudin - leech (Hirudinaria manillensis)
C:Species: Hirudinaria manillensis
C:Date: 27-Sep-1994 #sequence_revision 18-Nov-1994 #text_change 09-Mar-1996
C:Accession: A53883
R:Electrowala, A.; Hartwell, R.; Scaven, M.D.; Atkinson, T.
J. Protein Chem. 12, 365-370, 1993
A:Title: The complete amino acid sequence of a hirudin variant from the leech Hirudin
A:Reference number: A53883; MUID:94000343; PMID:8397794
A:Accession: A53883
A:Status: preliminary
A:Molecule type: protein
A:Residues: 1-63 <ELB>
A:Experimental source: head portions
A:Note: sequence extracted from NCBI backbone (NCBIP:139162)
C:Superfamily: thrombin inhibitor

Query Match 67.7%; Score 249; DB 2; Length 63;
Best Local Similarity 67.7%; Pred. No. 1.3e-18;
Matches 44; Conservative 8; Mismatches 11; Indels 2; Gaps 1;

QY 1 LVTDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 60
:|||||
DB 1 VYVDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 58

QY 61 EYIQQ 65
:|:
DB 59 DEXIK 63

RESULT 15

S33329
hirudin HM2 - leech (Hirudinaria manillensis)
C:Species: Hirudinaria manillensis
C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 16-Jul-1999
C:Accession: S33329
R:Scacheri, E.; Nitti, G.; Valsasina, B.; Orsini, G.; Visco, C.; Ferrera, M.; Sawyer,
Eur. J. Biochem. 214, 295-304, 1993
A:Title: Novel hirudin variants from the leech Hirudinaria manillensis. Amino acid se
A:Reference number: S33328; MUID:93285156; PMID:7685281
A:Accession: S33329
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-84 <SCA>
A:Cross-references: EMBL:X72786; NID:g313674; PIDN:CAA51293.1; PID:g313675
C:Genetics:
A:Introns: 21/1; 37/3; 61/1
C:Superfamily: thrombin inhibitor

Query Match 67.1%; Score 247; DB 2; Length 84;
Best Local Similarity 71.0%; Pred. No. 2.7e-18;
Matches 44; Conservative 6; Mismatches 10; Indels 2; Gaps 1;

QY 1 LVTDTCTESGNNLCICGSSNVCQGKNCILGSDGKNCQVTEGTPKPSHNDGDFEIP 60
:|||||
DB 21 VYVDTCTESGNNLCICGSSNVCQGKNCQVTEGTPKPSHNDGDFEIP 78

QY 61 EE 62
:|:
DB 79 DE 80

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Search completed: December 30, 2002, 16:18:00
Job time : 18 secs
